

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.

Application Serial Number: 09/462,929
Source: 1FW/b
Date Processed by STIC: 10/10/06

ENTERED



IFW16

RAW SEQUENCE LISTING

DATE: 10/10/2006

PATENT APPLICATION: US/09/462,929

TIME: 15:08:38

Input Set : N:\AMC\462929.txt

Output Set: N:\CRF4\10102006\I462929.raw

3 <110> APPLICANT: Old, Lloyd J.
 4 Scanlan, Matthew J.
 5 Stockert, Elisabeth
 6 Gure, Ali
 7 Chen, Yao-Tseng
 8 Gout, Ivan
 9 O'Hare, Michael
 10 Obata, Yuichi
 11 Pfreundschuh, Michael
 12 Tureci, Ozlem
 13 Sahin, Ugur
 16 <120> TITLE OF INVENTION: CANCER-ASSOCIATED NUCLEIC ACIDS AND
 17 POLYPEPTIDES
 19 <130> FILE REFERENCE: L0461.70078US00
 21 <140> CURRENT APPLICATION NUMBER: 09/462,929
 C--> 22 <141> **CURRENT FILING DATE: 2000-06-07**
 24 <150> PRIOR APPLICATION NUMBER: PCT/US98/14679
 25 <151> PRIOR FILING DATE: 1998-07-15
 27 <150> PRIOR APPLICATION NUMBER: US 08/896,164
 28 <151> PRIOR FILING DATE: 1997-07-17
 30 <150> PRIOR APPLICATION NUMBER: US 09/948,705
 31 <151> PRIOR FILING DATE: 1997-10-10
 33 <150> PRIOR APPLICATION NUMBER: US 09/102,322
 34 <151> PRIOR FILING DATE: 1998-06-22
 36 <150> PRIOR APPLICATION NUMBER: US 60/061,599
 37 <151> PRIOR FILING DATE: 1997-10-10
 39 <150> PRIOR APPLICATION NUMBER: US 60/061,765
 40 <151> PRIOR FILING DATE: 1997-10-10
 42 <150> PRIOR APPLICATION NUMBER: UK 9721697.2
 43 <151> PRIOR FILING DATE: 1997-10-11
 45 <160> NUMBER OF SEQ ID NOS: 822
 47 <170> SOFTWARE: PatentIn version 3.2
 49 <210> SEQ ID NO: 1
 50 <211> LENGTH: 474
 51 <212> TYPE: DNA
 52 <213> ORGANISM: Homo sapiens
 55 <220> FEATURE:
 56 <221> NAME/KEY: misc_feature
 57 <222> LOCATION: (26)..(26)
 58 <223> OTHER INFORMATION: n is a, c, g, or t
 60 <220> FEATURE:
 61 <221> NAME/KEY: misc_feature
 62 <222> LOCATION: (68)..(68)

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63 <223> OTHER INFORMATION: n is a, c, g, or t
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67 <222> LOCATION: (119)..(119) /
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72 <222> LOCATION: (138)..(138) /
73 <223> OTHER INFORMATION: n is a, c, g, or t
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76 <221> NAME/KEY: misc_feature
77 <222> LOCATION: (195)..(195) /
78 <223> OTHER INFORMATION: n is a, c, g, or t
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81 <221> NAME/KEY: misc_feature
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86 <221> NAME/KEY: misc_feature
87 <222> LOCATION: (265)..(265) /
88 <223> OTHER INFORMATION: n is a, c, g, or t
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92 <222> LOCATION: (274)..(274) /
93 <223> OTHER INFORMATION: n is a, c, g, or t
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121 <221> NAME/KEY: misc_feature
122 <222> LOCATION: (403)..(403) /
123 <223> OTHER INFORMATION: n is a, c, g, or t

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127 <222> LOCATION: (413)..(413) ✓
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132 <222> LOCATION: (417)..(417) ✓
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163 <223> OTHER INFORMATION: n is a, c, g, or t
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183 <223> OTHER INFORMATION: n is a, c, g, or t
185 <220> FEATURE:

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196 <221> NAME/KEY: misc_feature
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203 <223> OTHER INFORMATION: n is a, c, g, or t
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208 ctctggantc tggtcttcac acagagctgg gctcgagttt tcagtgcctt tgaatccang      120
210 cgctctgtct ctgggacngc ttcattcaag tcttgactta cgccatcttg gcaagatgtc      180
212 tgcttgccat ccaanttctc catggtatcc cctttanaga cttcaccaag agatgttctt      240
214 ctctcaaaga tggtgttatt attgntaact tgantcccat tttgcttcag taagcctgaa      300
216 tacctcagct tccaaatctc taatccgcac ttggaggcct tgcacttcca naagangngc      360
218 tttttcaaat cttcnagttt cnttcttctg ctctgtctt tgnatagatc aanctcnttt      420
220 tgntgnnagt cntntntnaa cntcnttgcg tgntggngng tcntttncan nggc          474
223 <210> SEQ ID NO: 2
224 <211> LENGTH: 1054
225 <212> TYPE: DNA
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251 <222> LOCATION: (85)..(86)
252 <223> OTHER INFORMATION: n is a, c, g, or t
254 <220> FEATURE:

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255 <221> NAME/KEY: misc_feature
256 <222> LOCATION: (89)..(89) ✓
257 <223> OTHER INFORMATION: n is a, c, g, or t
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260 <221> NAME/KEY: misc_feature
261 <222> LOCATION: (101)..(101) ✓
262 <223> OTHER INFORMATION: n is a, c, g, or t
264 <220> FEATURE:
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287 <223> OTHER INFORMATION: n is a, c, g, or t
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292 <223> OTHER INFORMATION: n is a, c, g, or t
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296 <222> LOCATION: (166)..(166) ✓
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309 <220> FEATURE:
310 <221> NAME/KEY: misc_feature
311 <222> LOCATION: (204)..(204) ✓
312 <223> OTHER INFORMATION: n is a, c, g, or t
314 <220> FEATURE:
315 <221> NAME/KEY: misc_feature

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/462,929

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 TIME: 15:08:39

FYI

Input Set : N:\AMC\462929.txt
 Output Set: N:\CRF4\10102006\I462929.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 26,68,119,138,195,217,265,274,351,356,358,375,382,403,413
 Seq#:1; N Pos. 417,423,426,427,433,435,436,438,442,445,450,453,457,458,460
 Seq#:1; N Pos. 463,467,470,471
 Seq#:2; N Pos. 27,29,30,46,58,85,86,89,101,103,107,110,128,155,157,166,170
 Seq#:2; N Pos. 200,204,208,225,227,234,236,238,244,255,258,268,271,276,278
 Seq#:2; N Pos. 292,300,307,309,317,323,330,331,335,343,350,356,363,366,375
 Seq#:2; N Pos. 377,378,413,425,430,431,437,439,444,449,462,463,466,469,482
 Seq#:2; N Pos. 499,500,506,508,509,510,513,523,529,531,532,535,541,549,570
 Seq#:2; N Pos. 576,596,600,607,614,615,616,619,622,624,630,636,638,641,643
 Seq#:2; N Pos. 644,654,655,659,663,670,676,693,694,698,700,702,703,708,721
 Seq#:2; N Pos. 722,724,727,729,731,741,748,751,752,753,754,756,758,760,764
 Seq#:2; N Pos. 767,768,774,780,782,788,789,799,806,808,817,818,820,821,826
 Seq#:2; N Pos. 827,839,841,848,849,860,862,863,864,870,871,874,882,886,888
 Seq#:2; N Pos. 896,898,900,902,929,930,952,959,964,966,968,973,980,981,985
 Seq#:2; N Pos. 1017,1025,1027,1033,1035,1039,1041,1047,1052
 Seq#:3; N Pos. 263,302,330,331,355,370,394,405,419,430,435
 Seq#:4; N Pos. 108,112,138,169,177,206,225
 Seq#:5; N Pos. 383,402,412
 Seq#:6; N Pos. 730
 Seq#:8; N Pos. 535,589,594,630,631,710
 Seq#:9; N Pos. 126,128,570
 Seq#:10; N Pos. 56,66,94,124,136,140,143,145,253,257,263,292,295,352,376
 Seq#:10; N Pos. 384,391,422,424,450,461,463,474,477,478,486,495,501,521,528
 Seq#:10; N Pos. 544,547,549,560,579,587,593,598,599,607,647,648,650,659
 Seq#:11; N Pos. 39,109,143,207,233,268,270,295,299,304,310,328,350,362,370
 Seq#:11; N Pos. 372,398,425,451,484,505,509,512,520,550,559,595,598,604,633
 Seq#:11; N Pos. 682,684,686,732,735,739,746,749,760,768,794,795,804,806,815
 Seq#:11; N Pos. 818,830,832,838,839,846,849,888,891,892,916,918,935,937,945
 Seq#:11; N Pos. 960,969,973,975,976,977,983,996,997,1000,1007,1011,1012
 Seq#:11; N Pos. 1019,1022,1023,1027,1030,1047,1049,1053,1061,1066,1067,1070
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 Seq#:12; N Pos. 307,546,549,577,587,597,600,616,645,673,674,684,686,687,693
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 Seq#:13; N Pos. 11,33,292,366,369
 Seq#:14; N Pos. 11,222,420,511,536,556,561,647,651,656,657,679,683,684,686
 Seq#:14; N Pos. 688,692,701,714,717,720,725,726,741,747,758,767,800,805,809
 Seq#:14; N Pos. 810,818,823,827,844,846,855,863,870,872,878,881,884,887,888
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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/462,929

DATE: 10/10/2006
TIME: 15:08:39

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Output Set: N:\CRF4\10102006\I462929.raw

Seq#:14; N Pos. 1121,1123,1125,1132,1133,1135,1139,1141,1145,1148,1149,1152
Seq#:14; N Pos. 1153,1156,1158,1159,1160,1162,1164
Seq#:15; N Pos. 34,312,451,530,542,566,570,627,628,629,632,638,646,650,653
Seq#:15; N Pos. 654,657,662,665,670,674,678,681,685,687,688,689,692,694,696
Seq#:15; N Pos. 700,704,711,715,716,717,718,719,723,732,741,745,748,749,753
Seq#:15; N Pos. 756,757,763,778,784,786,788,806,807,813,815,816,817,818,819
Seq#:15; N Pos. 822,823,827,828,841,842,844,846,849,850,851,852,854,855,856

Invalid <213> Response:

Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:817

VERIFICATION SUMMARY

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TIME: 15:08:39

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L:22 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:206 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
M:341 Repeated in SeqNo=1
L:955 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0
M:341 Repeated in SeqNo=2
L:1057 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:240
M:341 Repeated in SeqNo=3
L:1110 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:60
M:341 Repeated in SeqNo=4
L:1153 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:360
L:1196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:720
L:1274 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8 after pos.:480
M:341 Repeated in SeqNo=8
L:1311 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:120
M:341 Repeated in SeqNo=9
L:1540 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10 after pos.:0
M:341 Repeated in SeqNo=10
L:2017 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11 after pos.:0
M:341 Repeated in SeqNo=11
L:2260 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:300
M:341 Repeated in SeqNo=12
L:2313 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13 after pos.:0
M:341 Repeated in SeqNo=13
L:2795 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14 after pos.:0
M:341 Repeated in SeqNo=14
L:3383 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
M:341 Repeated in SeqNo=15
L:3773 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16 after pos.:60
M:341 Repeated in SeqNo=16
L:4009 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:60
M:341 Repeated in SeqNo=17
L:4399 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:540
M:341 Repeated in SeqNo=18
L:4594 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19 after pos.:0
M:341 Repeated in SeqNo=19
L:4701 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20 after pos.:180
M:341 Repeated in SeqNo=20
L:4752 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:180
M:341 Repeated in SeqNo=21
L:4895 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22 after pos.:120
M:341 Repeated in SeqNo=22
L:4971 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23 after pos.:180
M:341 Repeated in SeqNo=23
L:5028 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
M:341 Repeated in SeqNo=24
L:5094 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:120
M:341 Repeated in SeqNo=25
L:5223 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26 after pos.:0

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M:341 Repeated in SeqNo=26

L:5269 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:420

M:341 Repeated in SeqNo=27

L:65988 M:283 W: Missing Blank Line separator, <400> field identifier